

Ligand-receptor networks

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 An abbreviated version of this protocol was published in eLIFE in Mar 2019

Single-cell expression profiling reveals dynamic flux of cardiac stromal, vascular and immune cells in health and injury

DOI: 10.7554/eLife.43882

Detailed protocol

The method is implemented as an R package available at <https://github.com/VCCRI/scTalk>

How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Patrick, R. (2020). Ligand-receptor networks. Bio-protocol Preprint. [bio-protocol.org/prep309](https://doi.org/10.21203/rs.3.rs-333333/v1).
2. Farbehi, N., Patrick, R., Dorison, A., Xaymardan, M., Janbandhu, V., Wystub-Lis, K., Ho, J. W., Nordon, R. E. and Harvey, R. P. (2019). Single-cell expression profiling reveals dynamic flux of cardiac stromal, vascular and immune cells in health and injury. eLIFE. DOI: [10.7554/eLife.43882](https://doi.org/10.7554/eLife.43882)

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